CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/629,042E

1647 1600

CRF Processing Date: 12/19/02
Edited by: DC
Verified by: DC (STIC staff)

- [ ] Changed a file from non-ASCII to ASCII
- [ ] Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- [ ] Edited a format error in the Current Application Data section, specifically:

  ENTERED #39

  _____

- [ ] Edited the Current Application Data section with the actual current number. The number inputted by the applicant was [ ] the prior application data; or [ ] other _____.
- [ ] Added the mandatory heading and subheadings for "Current Application Data".
- [ ] Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- [ ] Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- [ ] Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- [ ] Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- [ ] Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- [ ] Inserted colons after headings/subheadings. Headings edited included: _____
- [ ] Deleted extra, invalid, headings used by an applicant, specifically: _____
- [✓] Deleted: [✓] non-ASCII "garbage" at the beginning/end of files; [ ] secretary initials/filename at end of file; [ ] page numbers throughout text; [ ] other invalid text, such as _____ .
- [ ] Inserted mandatory headings, specifically: _____
- [ ] Corrected an obvious error in the response, specifically: _____

  RECEIVED

  DEC 2 0 2002

  TECH CENTER 1600/2900

- [ ] Edited identifiers where upper case is used but lower case is required, or vice versa.
- [ ] Corrected an error in the Number of Sequences field, specifically: _____
- [ ] A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- [ ] Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- [ ] Other: _____

  _____
  _____
  _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

|||||||||||||||||||||||||||||||

1600

RAW SEQUENCE LISTING   DATE: 12/19/2002
PATENT APPLICATION: **US/09/029,042E**   TIME: 09:33:23

Input Set : **A:\PTO.DC.txt**
Output Set: **N:\CRF4\12192002\I029042E.raw**

```
 4 <110> APPLICANT: KIM, Sun-Young
 5        KIM, Kee-Won
 6        KIM, Tae-Han
 7        HWANG, Jeong-Ho
 8        KIM, Seon-Hee
 9        LEE, Sun-Young
11 <120> TITLE OF INVENTION: Heterologous Protein Production System Using Avian Cells
13 <130> FILE REFERENCE: 003364.P001
15 <140> CURRENT APPLICATION NUMBER: US 09/029,042E
16 <141> CURRENT FILING DATE: 1998-05-15
18 <150> PRIOR APPLICATION NUMBER: KR 10-1995-26391
19 <151> PRIOR FILING DATE: 1995-08-24
21 <150> PRIOR APPLICATION NUMBER: PCT/KR96/00145
22 <151> PRIOR FILING DATE: 1996-08-23
24 <160> NUMBER OF SEQ ID NOS: 15
26 <170> SOFTWARE: KopatentIn 1.71
29 <210> SEQ ID NO: 1
30 <211> LENGTH: 1585
31 <212> TYPE: DNA
32 <213> ORGANISM: Homo sapiens
34 <400> SEQUENCE: 1
35 atggggggtgc acgaatgtcc tgcctggctg tggcttctcc tgtccctgct gtcgctccct          60
36 ctgggcctcc cagtcctggg cgccccacca cgcctcatct gtgacagccg agtcctggag         120
37 aggtacctct tggaggccaa ggaggccgag aatatcacgg tgagacccct tccccagcac         180
38 attccacaga actcacgctc agggcttcag ggaactcctc ccagatccag gaacctggca         240
39 cttggttttgg ggtggagttg ggaagctaga cactgccccc ctacataaga ataagtctgg         300
40 tggccccaaa ccatacctgg aaactaggca aggagcaaag ccagcagatc ctacggcctg         360
41 tgggccaggg ccagagcctt cagggaccct tgactccccg ggctgtgtgc atttcagacg         420
42 ggctgtgctg aacactgcag cttgaatgag aatatcactg tcccagacac caaagttaat         480
43 ttctatgcct ggaagaggat ggaggtgagt tcctttttt ttttttttcc tttcttttgg         540
44 agaatctcat ttgcgagcct gattttggat gaaagggaga atgatcgggg gaaaggtaaa         600
45 atggagcagc agagatgagg ctgcctgggc gcagaggctc acgtctataa tcccaggctg         660
46 agatggccga gatggagaa ttgcttgagc cctggaggtt cagaccaacc taggcagcat         720
47 agtgagatcc cccatctcta caaacattta aaaaaattag tcaggtgaag tggtgcatgg         780
48 tggtagtccc agatatttgg aaggctgagg cgggaggatc gcttgagccc aggaatttga         840
49 ggctgcagtg agctgtgatc acaccactgc actccagcct cagtgacaga gtgaggccct         900
50 gtctcaaaaa agaaaagaaa aagaaaaat aatgagggct gtatggaata cattcattat         960
51 tcattcactc actcactcac tcattcattc attcattcat tcaacaagtc ttattgcata        1020
52 ccttctgttt gctcagcttg gtgcttgggg ctgctgaggg gcaggaggga gagggtgaca        1080
53 tgggtcagct gactcccaga gtccactccc tgtaggtcgg gcagcaggcc gtagaagtct        1140
54 ggcagggcct ggccctgctg tcggaagctg tcctgcgggg ccaggccctg ttggtcaact        1200
55 cttcccagcc gtgggagccc ctgcagctgc atgtggataa agccgtcagt ggccttcgca        1260
56 gcctcaccac tctgcttcgg gctctgggag cccaggtgag taggagcgga cacttctgct        1320
```

```
 57 tgccctttct gtaagaaggg gagaagggtc ttgctaagga gtacaggaac tgtccgtatt        1380
 58 ccttcccttt ctgtggcact gcagcgacct cctgtttttct ccttggcaga aggaagccat       1440
 59 ctcccctcca gatgcggcct cagctgctcc actccgaaca atcactgctg acactttccg         1500
 60 caaactcttc cgagtctact ccaatttcct ccggggaaag ctgaagctgt acacagggga         1560
 61 ggcctgcagg acaggggaca gatga                                               1585
 64 <210> SEQ ID NO: 2
 65 <211> LENGTH: 1583
 66 <212> TYPE: DNA
 67 <213> ORGANISM: Homo sapiens
 69 <400> SEQUENCE: 2
 70 atggggggtgc acgaatgtcc tgcctggctg tggcttctcc tgtccctgct gtcgctccct          60
 71 ctgggcctcc cagtcctggg cgccccacca cgcctcatct gtgacagccg agtcctggag         120
 72 aggtacctct tggaggccaa ggaggccgag aatatcacgg tgagacccct tccccagcac         180
 73 attccacaga actcacgctc agggcttcag gcaactcctc ccagatccag gaacctggca         240
 74 cttggtttgg ggtggagttg ggaagctaga cactgcccccc ctacataaga ataagtctgg        300
 75 tggccccaaa ccatacctgg aaactaggca aggagcaaag ccagcagatc ctacgcctgt         360
 76 ggccagggcc agagccttca gggacccttg actccccggg ctgtgtgcat ttcagacggg         420
 77 ctgtgctgaa cactgcagct tgaatgagaa tatcactgtc ccagacacca aagttaattt         480
 78 ctatgcctgg aagaggatgg aggtgagttc cttttttttt tttttcctt tcttttggag          540
 79 aatctcattt gcgagcctga ttttggatga aagggagaat gatcgaggga aaggtaaaat         600
 80 ggagcagcag agatgaggct gcctgggcgc agaggctcac gtctataatc ccaggctgag         660
 81 atggccgaga tgggagaatt gcttgagccc tggaggttca gaccaaccta ggcagcatag        720
 82 tgagatcccc catctctaca aacatttaaa aaaattagtc aggtgaagtg gtgcatggtg         780
 83 gtagtccag atatttggaa ggctgaggcg ggaggatcgc ttgagcccag gaatttgagg          840
 84 ctgcagtgag ctgtgatcac accactgcac tccagcctca gtgacagagt gaggccctgt         900
 85 ctcaaaaaag aaaagaaaaa agaaaaataa tgagggctgt atggaatacg ttcattattc         960
 86 attcactcac tcactcactc attcattcat tcattcattc aacatgtctt attgcatacc        1020
 87 ttctgtttgc tcagcttggt gcttggggct gctgaggggc aggagggaga gggtgacatc       1080
 88 cctcagctga ctcccagagt ccactccctg taggtcgggc agcaggccgt agaagtctgg        1140
 89 cagggcctgg ccctgctgtc ggaagctgtc ctgcgggggcc aggccctgtt ggtcaactct       1200
 90 tcccagccgt gggagcccct gcagctgcat gtggataaag ccgtcagtgg ccttcgcagc        1260
 91 ctcaccactc tgcttcgggc tctgggagcc caggtgagta ggagcggaca cttctgcttg        1320
 92 ccctttctgt aagaagggga gaagggtctt gctaaggagt acaggaactg tccgtattcc        1380
 93 ttcccttttct gtggcactgc agcgacctcc tgttttctcc ttggcagaag gaagccatct      1440
 94 ccccctccaga tgcggcctca gctgctccac tccgaacaat cactgctgac actttccgca       1500
 95 aactcttccg agtctactcc aatttcctcc ggggaaagct gaagctgtac acagggggag        1560
 96 cctgcaggac agggggacaga tga                                               1583
 99 <210> SEQ ID NO: 3
100 <211> LENGTH: 1585
101 <212> TYPE: DNA
102 <213> ORGANISM: Homo sapiens
104 <400> SEQUENCE: 3
105 atggggggtgc acgaatgtcc tgcctggctg tggcttctcc tgtccctgct gtcgctccct          60
106 ctgggcctcc cagtcctggg cgccccacca cgcctcatct gtgacagccg agtcctggag         120
107 aggtacctct tggaggccaa ggaggccgag aatatcacgg tgagacccct tccccagcac         180
108 attccacaga actcacgctc agggcttcag gcaactcctc ccagatccag gaacctggca         240
109 cttggtttgg ggtggagttg ggaagctaga cactgcccccc ctacataaga ataagtctgg        300
110 tggccccaaa ccatacctgg aaactaggca aggagcaaag ccagcagatc ctacggcctg         360
111 tgggccaggg ccaaaacctt cagggaccct tgactccccg ggctgtttgc atttcagacg        420
```

RAW SEQUENCE LISTING   DATE: 12/19/2002
PATENT APPLICATION: **US/09/029,042E**   TIME: 09:33:23

Input Set : **A:\PTO.DC.txt**
Output Set: **N:\CRF4\12192002\I029042E.raw**

```
112 ggctgtgctg aacactgcag cttgaatgaa aatatcactg tcccagacac caaagttaat      480
113 ttctatgcct ggaagaggat ggaggtgagt tccttttttt tttttttttcc tttctttttgg      540
114 agaatctcat ttgcgagcct gattttggat gaaagggaga atgatcgagg gaaaggtaaa      600
115 atggagcagc agagatgagg ctgcctgggc gcagaggctc acgtctataa tcccaggctg      660
116 agatggccga aatgggagaa ttgcttgagc cctggaggtt cagaccaacc taggcagcat      720
117 agtgagatcc cccatctcta caaacattta aaaaaattag tcaggtgaag tggtgcatgg      780
118 tggtagtccc agatatttgg atggctgagg cgggaggatc gcttgagccc aggaatttga      840
119 ggctgcagtg agctgtgatc acaccactgc actccagcct cagtgacaga atgaggccct      900
120 gtctcaaaaa agaaaagaaa aagaaaaat aatgagggct gtatggaata cattcattat      960
121 tcattcactc actcactcac tcatccattc attcattcat tcaacaagtc ttattgcata     1020
122 ccttctgttt gctcagcttg gtgctcgggg ctgctgaggg gcaggaggga gagggtgaca     1080
123 tgggtcagct gactcccaga gtccactccc tgtaggtcgg gcaacaggcc gtagaagtct     1140
124 ggcagggcct ggccctgctg tcggaagctg tcctgcgggg ccaggccctg ttggtcaact     1200
125 tttcccagcc gtgggagccc ctgcagctgc atgtggataa agccgtcagt ggccttcgca     1260
126 gcctcaccac tctgcttcgg gctctgggag cccaggtgag taggagcgga cacttctgct     1320
127 tgccctttct gtaagaaggg gagaagggtc ttgctaagga gtacaggaac tgtccgtatt     1380
128 ccttcccttt ctgtggcact gcagcgacct cctgttttct ccttggcaga aggaagccat     1440
129 ctcccctcca gatgcggcct cagctgctcc actccgaaca atcactgctg acactttccg     1500
130 caaactcttc cgagtctact ccaatttcct ccggggaaag ctgaagctgt acacagggga     1560
131 ggcctgcagg acagggggaca gatga                                          1585
134 <210> SEQ ID NO: 4
135 <211> LENGTH: 1586
136 <212> TYPE: DNA
137 <213> ORGANISM: Homo sapiens
139 <400> SEQUENCE: 4
140 atggggtgc acgaatgtcc tgcctggctg tggcttctcc tgtccctgct gtcgctccct       60
141 ctgggcctcc cagtcctggg cgccccacca cgcctcatct gtgacagccg agtcctggag      120
142 aggtacctct tggaggccaa ggaggccgag aatatcacgg tgagacccct tccccagcac      180
143 attccacaga actcacgctc agggcttcag gcaactcctc ccagatccag gaacctggca      240
144 cttggtttgg ggtggagttg ggaagctaga cactgccccc ctacataaga ataagtctgg      300
145 tggccccaaa ccatacctgg aaactaggca aggagcaaag ccagcagatc ctacggcctg      360
146 tgggccaggg ccaggagcct tcaggaccc ttgactcccc gggctgtgtg catttcagaa      420
147 gggctgtgct gaacactgca gcttgaatga gaatatcact gtcccagaca ccaaagttaa      480
148 tttctatgcc tggaagagga tggaggtgag ttcctttttt ttttttttttc ctttctttttg      540
149 gagaatctca tttgcgagcc tgattttgga tgaaagggag agtgatcgag ggaaaggtaa      600
150 aatggagcag cagagatgag gctgcctggg cgcagaggct cacgtctata atcccaggct      660
151 gagatggccg agatgggaga attgcttgag ccctggaggt tcagaccaac ctaggcagca      720
152 tagtgagatc ccccatctct acaaacattt aaaaaaatta gtcaggtgaa gtggtgcatg      780
153 gtggtagtcc cagatatttg gaaggctgag gcgggaggat cgcttgagcc caggaatttg      840
154 aggctgcggt gagctgtgat cacaccactg cactccagcc tcagtgacag agtgagccc      900
155 tgtctcaaaa aagaaaagaa aaaagaaaaa taatgagggc tgtatggaat acattcatta      960
156 ttcattcact cactcactca ctcattcat cattcattca ttcaacaagt cttattgcat     1020
157 accttctgtt tgctcagctt ggtgcttggg gctgctgagg ggcaggaggg tgagggtgac     1080
158 atgggtcagc tgactcccag agtccactcc ctgttggtcg ggcagcaggc cgtagaagtc     1140
159 tggcagggcc tggccctgct gtcggaagct gtcctgcggg gccaggccct gttggtcaac     1200
160 tcttcccagc cgtgggagcc cctgcagctg catgtggata aagccgtcag tggccttcgc     1260
161 agcctcacca ctctgcttcg ggctctggga gcccaggtga gtaggagcgg acacttctgc     1320
162 ttgccctttc gtaagaagg ggagaagggt cttgctaagg agtacaggaa ctgtccgtat     1380
163 tccttccctt tctgtggcac tgcagcgacc tcctgttttc tccttggcag aaggaagcca     1440
```

```
164 tctcccctcc agatgcggcc tcagctgctc cactccgaac aatcactgct gacactttcc          1500
165 gcaaactctt ccgagtctac tccaatttcc tccggggaaa gctgaagctg tacacagggg          1560
166 aggcctgcag gacaggggac agatga                                               1586
169 <210> SEQ ID NO: 5
170 <211> LENGTH: 1583
171 <212> TYPE: DNA
172 <213> ORGANISM: Homo sapiens
174 <400> SEQUENCE: 5
175 atggggstgc acgaatgtcc tgcctggctg tggcttctcc tgtccctgct gtcgctccct            60
176 ctgggcctcc cagtcctggg cgccccacca cgcctcatct gtgacagacg agtcctggag           120
177 aggtacctct tggaggccaa ggaggccgag aatatcacgg tgagacccct tccccagcac           180
178 attccacaga actcacgctc agggcttcag gcaactcctc ccagatccag gaacctggca           240
179 cttggtttgg ggtggagttg ggaagctaga cactgccccc ctacataaga ataagtctgg           300
180 tggccccaaa ccatacctgg aaactaggca aggagcaaag ccagcagatc ctacggcctg           360
181 tgggccaggg gcagagcctt cagggaccct tgactccccg ggctgtgtgc atttcagacg           420
182 ggctgtgctg aacactgcag cttgaatgag aatatcactg tcccagacac caaagttaat          480
183 ttctatgcct ggaagaggat ggaggtgagt tccttttttt tttttttcctt tcttttggag          540
184 aatctcattt gcgagcctga ttttggatga aagggagaat gatcgaggga aggtaaaat            600
185 ggagcagcag agatgaggct gcctgggcgc agaggctcac gtctataatc ccaggctgag           660
186 acggccgaga tgggagaatt gcttgagccc tggaggttca gaccaaccta ggcagcatag           720
187 tgagatcccc catctctaca aacatttaaa aaaattagtc aggtgaagtg gtgcatggtg           780
188 gtagtcccag atatttggaa ggctgaggcg ggaggatcgc ttgagcccag gaatttgggg           840
189 ctgcagtgag ctgtgatcac accactgcaa tccagcctca gtgacagagt gaggccctgt           900
190 ctcaaaaacg aaaagaaaaa agaaaaataa tgagggctgt atggaataca ttcattattc           960
191 attcactcac tcactcactc attcattcat tcattcattc aacaagtctt attgcatacc          1020
192 ttctgtttgc tcagcttggt gcttgggcct tctgaggggc aggagggaga gggtgacatg          1080
193 ggtcagctga ctcccagagt ccactccctg taggtcgggc agcaggccgt agaagtctgg          1140
194 cagggcctgg ccctgctgtc ggaatctgtc ctgcggggcc aggccctgtt ggtcaactct          1200
195 tcccaaccgt gggagcccct gcagctgcat gtggataaag ccgtcagtgg ccttcgcagc          1260
196 ctcaccactc tgcttcgggc tctgggagcc cagttgagta ggaggggaca cttctgcttg          1320
197 ccctttgtgt aagaaggaga gaagggtctt gctaaggagt acaggaactg tccgtattcc          1380
198 ttcccttttct gtggcactgc agcgacctcc tgttttctcc ttggcagaag gaagccatct         1440
199 cccctccaga tgcggcctca gctgctccac tccgaacaat cactgctgat actttccgca          1500
200 aactcttccg agtctactcc aatttcctcc ggggaaagct gaagctgtac acaggggagg          1560
201 cctgcaggac aggggacaga tga                                                  1583
204 <210> SEQ ID NO: 6
205 <211> LENGTH: 1587
206 <212> TYPE: DNA
207 <213> ORGANISM: Homo sapiens
209 <400> SEQUENCE: 6
210 atggggstgc acgaatgtcc tgcctggctg tggcttctcc tgtccctgct gtcgctccct            60
211 ctgggcctcc cagtcctggg cgccccacca cgcctcatct gtgacagccg agtcctggag           120
212 aggtacctct tggaggccaa ggaggccgag aatatcacgg tgagacccct tccccagcac           180
213 attccacaga actcacgctc agggcttcag gcgaactcct cccaggatcc aggaacctgg           240
214 cacttggttt ggggtggagt tgggaagcta gacactgccc ccctacataa gaataagtct           300
215 ggtggcccca aaccatacct ggaaactagg caaggagcaa agccagcaga tcctacggcc           360
216 tgtgggccag ggccagagcc ttcagggacc cttgactccc cgggctgtgt gcattccaga           420
217 cgggctgtgc tgaacactgc agcttgaatg agaatatcac tgtcccagac accaaagtta           480
218 atttctatgc ctggaagagg atggaggtga gttccttttt tttttttttt cctttctttt           540
```

Input Set : **A:\PTO.DC.txt**
Output Set: **N:\CRF4\12192002\I029042E.raw**

```
219 ggagaatctc atttgcgagc ctgatttggg atgaaaggga gaatgatcga gggaaaggta    600
220 aaatggagca gcagagatga ggctgcctgg gcgcagaggc tccagtctat aatcccaggc    660
221 tgagatggcc gagatgggag aattgcttga gccctggagg ttcagaccaa cctaggcagc    720
222 ctagtgagat cccccatctc tacaaacatt taaaaaaatt agtcaggtga agtggtgcat    780
223 ggtggtagtc ccagatattt ggaaggctga ggcgggagga tcgcttgagc ccaggaattt    840
224 gaggctgcag tgagctgtga tcacaccact gcactccagc ctcagtgaca gagtgaggcc    900
225 ctgtctcaaa aaagaaaaga aaaagaaaa attatgaggg ctgtatggaa tacattcatt     960
226 attcattcac tcactcactc actcattcat tcattcattc attcaacaag tcttattgca   1020
227 taccttctgt ttgctcagct tggtgcttgg ggctgctgag gggcaggagg gagagggtga   1080
228 catgggtcaa ctgactccca gagtccactc cctgtaggtc gggcagcagg ccgtagaagt   1140
229 ctggcagggc ctggccctgc tgtcggaagc tgtcctgcgg ggccaggccc tgttggtcaa   1200
230 ctcttcccag ccgtgggagc ccctgcagct gcatgtggat aaagccgtca gtggccttcg   1260
231 cagcctcacc actctgcttc gggctctggg agcccaggtg agtaggagcg gacacttctg   1320
232 cttgcccttt ctgtaagaag gggagaaggg tcttgctaag gagtacagga tctgtccgta   1380
233 ttccttccct ttctgtggca ctgcagcgac cacctgtttt ctccttggca gaaggaagcc   1440
234 atctcccctc cagatgcggc ctcagctgct ccactccgaa caatcactgc tgacactttc   1500
235 cgcaaactct tccgagtcta ctccaatttc ctccggggag agctgaagct gtacacaggg   1560
236 gaggcctgca ggacagggga cggatga                                       1587
239 <210> SEQ ID NO: 7
240 <211> LENGTH: 193
241 <212> TYPE: PRT
242 <213> ORGANISM: Homo sapiens
244 <400> SEQUENCE: 7
245 Met Gly Val His Glu Cys Pro Ala Trp Leu Trp Leu Leu Leu Ser Leu
246 1               5                   10                  15
247 Leu Ser Leu Pro Leu Gly Leu Pro Val Leu Gly Ala Pro Pro Arg Leu
248             20                  25                  30
249 Ile Cys Asp Ser Arg Val Leu Glu Arg Tyr Leu Leu Glu Ala Lys Glu
250         35                  40                  45
251 Ala Glu Asn Ile Thr Thr Gly Cys Ala Glu His Cys Ser Leu Asn Glu
252     50                  55                  60
253 Asn Ile Thr Val Pro Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg
254 65                  70                  75                  80
255 Met Glu Val Gly Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu
256             85                  90                  95
257 Leu Ser Glu Ala Val Leu Arg Gly Gln Ala Leu Leu Val Asn Ser Ser
258             100                 105                 110
259 Gln Pro Trp Glu Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly
260         115                 120                 125
261 Leu Arg Ser Leu Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu
262     130                 135                 140
263 Ala Ile Ser Pro Pro Asp Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile
264 145                 150                 155                 160
265 Thr Ala Asp Thr Phe Arg Lys Leu Phe Arg Val Val Ser Asn Phe Leu
266             165                 170                 175
267 Arg Gly Lys Leu Lys Leu Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp
268             180                 185                 190
269 Arg
272 <210> SEQ ID NO: 8
```

VERIFICATION SUMMARY   DATE: 12/19/2002
PATENT APPLICATION: **US/09/029,042E**   TIME: 09:33:24

Input Set : **A:\PTO.DC.txt**
Output Set: **N:\CRF4\12192002\I029042E.raw**